



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/728,420A
Source: 1 Fwd
Date Processed by STIC: 8/13/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

~~TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER~~
VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>) , EFS Submission User Manual - cPAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box-1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/728,420A

DATE: 08/13/2004

TIME: 16:13:54

Input Set : D:\Sequence Listing-rev.txt

Output Set: N:\CRF4\08132004\J728420A.raw

4 <110> APPLICANT: WOULFE, SUSAN L.
5 JAIN, RITA
6 BURR, AIMEE
8 <120> TITLE OF INVENTION: ENGINEERED FAB' FRAGMENT ANTI-TUMOR
9 NECROSIS FACTOR ALPHA IN COMBINATION WITH DISEASE MODIFYING
10 ANTI-RHEUMATIC DRUGS
12 <130> FILE REFERENCE: 122294-1010
14 <140> CURRENT APPLICATION NUMBER: US/10/728,420A
16 <141> CURRENT FILING DATE: 2003-12-05
18 <150> PRIOR APPLICATION NUMBER: US 60/431,053
20 <151> PRIOR FILING DATE: 2002-12-05
E--> 22 <160> NUMBER OF SEQ ID NOS: 40 (117 (P.2))
24 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

E--> 389 <210> SEQ ID NO: 46

(see p.3)

<210> 117
<211> 1477
<212> DNA
<213> Artificial Sequence

*last sequence in submitted
file*

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2

<220>

<223> Synthetic CDP870 nucleic acid sequence (anti-sense strand)

<400> 117

```
tcacacgcg gcgcattgtg gagttttgtc acaagatttg ggctcaactt tcttgtggac 60
cttggtgttg ctgggcttgt gattcacgtt gcagatgtag gtctgggtgc ccaagctgct 120
ggagggcacg gtcaccacgc tgctgagga gtagagtcct gaggactgta ggacagccgg 180
gaaggtgtgc acgcccgttg tcagggcgcc tgagttccac gacaccgtca ccggttcggg 240
gaagtagtcc ttgaccaggc agcccagggc cgctgtgccc ccagaggtgc tcttggagga 300
gggtgccagg gggaagaccg atgggccctt ggtggaagct gaggagactg tgactagggt 360
accctggccc cagtagtcca tggcataaga tctgtatcct ctacacaaat agtacactgc 420
ggtgtcctct gctctcaggc tattcatttg gaggtatgct gttgacttgg atgtgtctag 480
agagaacgtg aatctcgctt tgacgtgtc agcataaata ggctctccaa tgtaagtatt 540
aatccaaccc atccattcga ggccctttcc cggggcctgt ctaacccaat tcattccata 600
gtctgtgaag acgtaaccag atgcagcaca ggacaatctc agtgcacgc caggctgcac 660
gagaccgcct cctgactcga ccagctgaac ctcagcttgc gccacgggtg cgaaaccagc 720
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ctctattaaa gctttttgtt actggtgagc tcaggccctg atgggtgact tcgcaggcgt 840
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tgctgtcctt gctgtcctgc tctgtgacac tctcctggga gttaccgat tggaggcgt 960
tatccacctt ccactgtact ttggcctctc tgggataga gttattcagc aggcacacaa 1020
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ccgctaccgt acgtttgatt tctactttag taccctgacc gaatgtgagt gggtagatgt 1140
tatactgttg acagtaataa gtggcgaaat cttctggctg gaggctacta gtcgtgaggg 1200
tgaaatcagt accactaccg gatccgctga acctgtatgg tacaccacta tagaggaaag 1260
aggcactgta gatgagggct tttggggcct tacctgggtt ttgotgatac caggctacgt 1320
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tcaggctgga tgggctctgg gtcatttgaa tgtcagcttg cgctacggta gcgaaaccag 1440
ccaaggccac tgcaattgcg atagctgtct ttttcat 1477
```

replace with

\downarrow
 $\langle 400 \rangle$ $\langle 210 \rangle$ 46
000

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/728,420A

DATE: 08/13/2004

TIME: 16:13:55

Input Set : D:\Sequence Listing-rev.txt

Output Set: N:\CRF4\08132004\J728420A.raw

L:162 M:283 W: Missing Blank Line separator, <400> field identifier
L:163 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (9) SEQUENCE:
L:167 M:283 W: Missing Blank Line separator, <400> field identifier
L:168 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (10) SEQUENCE:
L:184 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:219 M:283 W: Missing Blank Line separator, <400> field identifier
L:220 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (12) SEQUENCE:
L:224 M:283 W: Missing Blank Line separator, <400> field identifier
L:225 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (13) SEQUENCE:
L:229 M:283 W: Missing Blank Line separator, <400> field identifier
L:230 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (14) SEQUENCE:
L:234 M:283 W: Missing Blank Line separator, <400> field identifier
L:235 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (15) SEQUENCE:
L:239 M:283 W: Missing Blank Line separator, <400> field identifier
L:240 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (16) SEQUENCE:
L:244 M:283 W: Missing Blank Line separator, <400> field identifier
L:245 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (17) SEQUENCE:
L:249 M:283 W: Missing Blank Line separator, <400> field identifier
L:250 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (18) SEQUENCE:
L:254 M:283 W: Missing Blank Line separator, <400> field identifier
L:255 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (19) SEQUENCE:
L:259 M:283 W: Missing Blank Line separator, <400> field identifier
L:260 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (20) SEQUENCE:
L:264 M:283 W: Missing Blank Line separator, <400> field identifier
L:265 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (21) SEQUENCE:
L:269 M:283 W: Missing Blank Line separator, <400> field identifier
L:270 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (22) SEQUENCE:
L:274 M:283 W: Missing Blank Line separator, <400> field identifier
L:275 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (23) SEQUENCE:
L:279 M:283 W: Missing Blank Line separator, <400> field identifier
L:280 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (24) SEQUENCE:
L:284 M:283 W: Missing Blank Line separator, <400> field identifier
L:285 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (25) SEQUENCE:
L:289 M:283 W: Missing Blank Line separator, <400> field identifier
L:290 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (26) SEQUENCE:
L:294 M:283 W: Missing Blank Line separator, <400> field identifier
L:295 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (27) SEQUENCE:
L:299 M:283 W: Missing Blank Line separator, <400> field identifier
L:300 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (28) SEQUENCE:
L:304 M:283 W: Missing Blank Line separator, <400> field identifier
L:305 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (29) SEQUENCE:
L:309 M:283 W: Missing Blank Line separator, <400> field identifier
L:310 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (30) SEQUENCE:
L:314 M:283 W: Missing Blank Line separator, <400> field identifier
L:315 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (31) SEQUENCE:
L:319 M:283 W: Missing Blank Line separator, <400> field identifier
L:320 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (32) SEQUENCE:
L:324 M:283 W: Missing Blank Line separator, <400> field identifier

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Input Set : D:\Sequence Listing-rev.txt

Output Set: N:\CRF4\08132004\J728420A.raw

L:325 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (33) SEQUENCE:
L:329 M:283 W: Missing Blank Line separator, <400> field identifier
L:330 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (34) SEQUENCE:
L:334 M:283 W: Missing Blank Line separator, <400> field identifier
L:335 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (35) SEQUENCE:
L:339 M:283 W: Missing Blank Line separator, <400> field identifier
L:340 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (36) SEQUENCE:
L:344 M:283 W: Missing Blank Line separator, <400> field identifier
L:345 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (37) SEQUENCE:
L:349 M:283 W: Missing Blank Line separator, <400> field identifier
L:350 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (38) SEQUENCE:
L:354 M:283 W: Missing Blank Line separator, <400> field identifier
L:355 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (39) SEQUENCE:
L:359 M:283 W: Missing Blank Line separator, <400> field identifier
L:360 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (40) SEQUENCE:
L:364 M:283 W: Missing Blank Line separator, <400> field identifier
L:365 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (41) SEQUENCE:
L:369 M:283 W: Missing Blank Line separator, <400> field identifier
L:370 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (42) SEQUENCE:
L:374 M:283 W: Missing Blank Line separator, <400> field identifier
L:375 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (43) SEQUENCE:
L:379 M:283 W: Missing Blank Line separator, <400> field identifier
L:380 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (44) SEQUENCE:
L:384 M:283 W: Missing Blank Line separator, <400> field identifier
L:385 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (45) SEQUENCE:
L:389 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO:46
L:389 M:283 W: Missing Blank Line separator, <210> field identifier
L:390 M:259 W: Allowed number of lines exceeded, <210> SEQ ID NO
L:394 M:283 W: Missing Blank Line separator, <400> field identifier
L:395 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (47) SEQUENCE:
L:399 M:283 W: Missing Blank Line separator, <400> field identifier
L:400 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (48) SEQUENCE:
L:404 M:283 W: Missing Blank Line separator, <400> field identifier
L:405 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (49) SEQUENCE:
L:409 M:283 W: Missing Blank Line separator, <400> field identifier
L:410 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (50) SEQUENCE:
L:414 M:283 W: Missing Blank Line separator, <400> field identifier
L:415 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (51) SEQUENCE:
L:419 M:283 W: Missing Blank Line separator, <400> field identifier
L:420 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (52) SEQUENCE:
L:424 M:283 W: Missing Blank Line separator, <400> field identifier
L:425 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (53) SEQUENCE:
L:429 M:283 W: Missing Blank Line separator, <400> field identifier
L:430 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (54) SEQUENCE:
L:434 M:283 W: Missing Blank Line separator, <400> field identifier
L:435 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (55) SEQUENCE:
L:439 M:283 W: Missing Blank Line separator, <400> field identifier
L:440 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (56) SEQUENCE:
L:444 M:283 W: Missing Blank Line separator, <400> field identifier

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Input Set : D:\Sequence Listing-rev.txt

Output Set: N:\CRF4\08132004\J728420A.raw

L:445 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (57) SEQUENCE:
L:449 M:283 W: Missing Blank Line separator, <400> field identifier
L:450 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (58) SEQUENCE:
L:454 M:283 W: Missing Blank Line separator, <400> field identifier
L:455 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (59) SEQUENCE:
L:460 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (60) SEQUENCE:
L:890 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:894 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:101
L:913 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:102
L:937 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:103
L:961 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:104
L:985 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:105
L:22 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (40) Counted (116)